Abade, A.

see Trovoada, M. J. et al.

ABO-secretor locus (FUT2). Polymorphism of the human ... in four populations in Asia: indication of distinct Asian subpopulations. 429

Adinolfi. M.

see Cirigliano, V. et al.

affected sibling pairs. Sample size requirements to control for stochastic variation in magnitude and location of allele-sharing linkage statistics in, 491

African prehistory. Phylogeography of the human mitochondrial haplogroup L3e: a snapshot of... and Atlantic slave trade, 549

Akar, N

see Malaspina. P. et al.

Alberman, E.

see Hermon, C. et al.

Alberman, E.

see Swerdlow, A. J. et al.

allele frequency and allele positivity. Comparison of statistical power between 2×2...tables in case control studies of complex disease genes, 197

allelic association discriminates draft orders, 503

Allende, L.

see Martinez-Laso, J. et al.

altitude adaptation. Angiotensin converting enzyme insertion allele in relation to high, 531

Aluja, M. P.

see Giraldo, M. P. et al.

Alves, C.

see Gusmao. L. et al.

Alves. C

see Trovoada, M. J. et al.

Alves-Silva, J.

see Bandelt, H. J. et al.

Alzheimer's disease. A novel method of two-locus linkage analysis applied to a genome scan for late onset, 473

Amerindians. An HLA class-II allele frequent in Eskimos and ... is found in the Tyrolean Ice Man, 363

Amorim, A

see Bandelt, H. J. et al.

Amorim. A.

see Gusmao. L. et al.

Amorim. A

see Pereira, L. et al.

Amorim, A.

see Trovoada, M. J. et al.

Anagnou, N

see Malaspina, P. et al.

Andersen-Ranberg, K.

see Varcasia. O. et al.

Andreev, K

see Varcasia, O. et al.

an euploidies. Assessment of new markers for the rapid detection of ... by quantitative fluorescent PCR (QF-PCR), 421

angiotensin converting enzyme insertion allele in relation to high altitude adaptation, 531

APOB-VNTR locus. Replication studies in longevity: puzzling findings in Danish centenarians at the 3'. apolipoprotein C-HL Interaction of the common... (APOC3 - 482C > T) and hepatic lipase (LIPC - 514C > T) promoter variants affects glucose tolerance in young adults. European Atherosclerosis Research Study II (EARS-II), 237

Arnaiz-Villena. A.

see Martinez-Laso, J. et al.

artificial neural network. Use of an...to detect association between a disease and multiple marker genotypes, 95

Asian populations. Distribution of FMR1 and FMR2 alleles in Javanese: confirmation of a specific AGGinterruption pattern in, 127

Asian subpopulations. Polymorphism of the human ABOsecretor locus (FUT2) in four populations in Asia: indication of distinct, 429

association. Sample size calculations for classical... and TDT-type methods using family data, 293

association. Family based tests of ... and/or linkage, 407 Atlantic. Phylogeography of the human mitochondrial haplogroup L3e: a snapshot of African prehistory and ... slave trade, 549

Babron, M. C.

see Greco. L. et al.

Baciliero, U

see Martinelli, M. et al.

Backes-Duro, Ch.

see Giraldo, M. P. et al.

Bandelt, H. J., Alves-Silva, J., Guimaraes, P. E. M., Santos, M. S., Brehm, A., Pereira, L., Coppa, A., Larruga, J. M., Rengo, C., Scozzari, R., Torroni, A., Prata, M. J., Amorim, A., Prado, V. F. & Pena, S. D. J.

phylogeography of the human mitochondrial haplogroup L3e: a snapshot of African prehistory and Atlantic slave trade, 549

Bantu expansions. Prehistoric and historic traces in the mtDNA of Mozambique: insights into the ... and the slave trade, 439

Barbujani. G.

see Malaspina. P. et al.

Barros, F

see Vega. A. et al.

Bathum, L.

see Varcasia, O, et al.

Benammar Elgaaied, A

see Khodjet El Khil, H. et al.

Beral, V.

see Hermon, C. et al.

Beral, V.

see Swerdlow, A. J. et al.

Beraud-Colomb. E.

see Mogentale-Profizi. N. et al.

Bertranpetit, J.

see Laluela-Fox. C. et al.

Bonafe, M.

see Varcasia, O. et al.

Brahmachari, S. K.

see Qadar Pasha, M. A. et al.

- Brehm, A.
 - see Bandelt, H. J. et al.
- Brewer, G. J.
 - see Olivarez. L. et al.
- Buckley, M. F.
 - see Faradz, M. H. et al.
- Calafell F
- see Laluela-Fox, C. et al.
- Calderon, F. L.
- see Laluela-Fox, C. et al.
- Camp, N. J. & Farnham, N. J.
 - correcting for multiple analyses in genomewide linkage studies, 577
- cancer incidence. Mortality and ... in persons with Down's syndrome, their parents and siblings, 167
- cancer incidence. Mortality and ... in persons with numerical sex chromosome abnormalities: a cohort study, 177
- Caribbean, MtDNA from extinct Tainos and the peopling of the, 137
- Carinci, F.
 - see Martinelli, M. et al.
- Carinci, P.
 - see Martinelli, M. et al.
- Carollo, C.
 - see Zapata, C. et al.
- Carracedo, A.
 - see Vega. A. et al.
- Casari G
 - see Greco. L. et al.
- case control studies. Comparison of statistical power between 2×2 allele frequency and allele positivity tables in ... of complex disease genes, 197
- Caucasian population. Estimate of the frequency of Wilson's disease in the U.S....: a mutation analysis approach, 459
- Caucasoid. HLA molecular markers in Tuvinians: a population with both Oriental and ... characteristics, 245
- Cavalli-Sforza, L. L.
 - see Underhill, P. A. et al.
- Cheng, R.
 - see Lin, S. et al.
- Chollet, L.
 - see Mogentale-Profizi, N. et al.
- chromosome 1. Linkage analysis of three candidate regions of ... in nonsyndromic familial orofacial eleft, 465
- chromosome 5q. Existence of a genetic risk factor on ... in Italian Coeliac Disease families, 35
- chromosome 11p11. Coeliac Disease: follow-up linkage study provides further support for existence of a susceptibility locus on, 377
- chromosome 14q24.3-q31. Grade-of-membership sibpair linkage analysis maps IDDM11 to, 387
- Ciaverella, G.
- see Malaspina, P. et al.
- Ciclitira, P. J.
 - see King, A. L. et al.
- Cirigliano, V., Lewin, P., Szpiro-Tapies, S., Fuster, C. &
 - assessment of new markers for the rapid detection of an euploidies by quantitative fluorescent PCR (QF-PCR), 421

- cleft lip and palate. Nonsyndromic . . . : complex genetics and environmental effects, 505
- Clerget-Darpoux, F.
- see Greco, L. et al.
- Clot. F
 - see Greco, L. et al.
- coeliac Disease. Existence of a genetic risk factor on chromosome 5q in Italian...families, 35
- coeliac Disease: follow-up linkage study provides further support for existence of a susceptibility locus on chromosome 11p11, 377
- Collins, A.
- see Ennis, S. et al.
- Coppa, A.
 - see Bandelt, H. J. et al.
- Corazza, G. R.
- see Greco, L. et al.
- Cordell, H. J.
 - sample size requirements to control for stochastic variation in magnitude and location of allele-sharing linkage statistics in affected sibling pairs, 491
- Corder, E. H., Woodbury, M. A., Manton, K. G. & Field, L. L.
- grade-of-membership sibpair linkage analysis maps *IDDM11* to chromosome 14q24.3-q31, 387
- Costas, J.
 - see Vega, A. et al.
- crossover interference. Genetic ... in the human genome, 79
- Curtis, D., North, B. V., & Sham, P. C.
- use of an artificial neural network to detect association between a disease and multiple marker genotypes, 95
- Curtis, D., North, B. V., & Sham, P. C.
- a novel method of two-locus linkage analysis applied to a genome scan for late onset Alzheimer's disease, 473 Curtis, D.
- see King, A. L. et al.
- D'Alfonso, A.
- see Greco, L. et al.
- D' measure, Sampling variance and distribution of the ... of overall gametic disequilibrium between multiallelic loci, 395
- Daker, M.
- see Swerdlow, A. J. et al.
- Danish. Replication studies in longevity: puzzling findings in ... centenarians at the 3' APOB-VNTR locus, 371
- Davis, M. B.
- see Vaughan, J. R. et al.
- De Benedictis, G.
- see Varcasia, O. et al.
- De Jonghe, P.
- see Irobi, J. et al.
- D. I.-.l.
- De Jonk, J. see Jira, P. E. et al.
- Delhanty, J. D. A.
 - preimplantation genetics: a review, 331
- Deng, H. W., Li, J. & Recker, R. R.
- LÕD score exclusion analyses for candidate genes using random population samples, 313
- Dearlove, A. M.
- see King. A. L. et al.
- Derenko, M. V.
- see Malyarchuk, B. A.

De Virgiliis, S.

see Greco, L. et al.

Dong, J. P.

see Jiang, R. et al.

Down's syndrome. Mortality and cancer incidence in persons with... their parents and siblings, 167

Dubut, V.

see Mogentale-Profizi. N. et al.

Dugoujon, J. M.

see Giraldo, M. P. et al.

Duman. T

see Malaspina, P. et al.

Ellis, H. J.

see King, A. L. et al.

Enhnolm, C

see Jansen. H. et al.

Ennis, S., Collins, A., Tapper, W., Murray, A., MacPherson, J. N. & Morton, N. E.

allelic association discriminates draft orders, 503 environmental effects. Nonsyndromic cleft lip and palate: Complex genetics and, 505

epistasis and its possible effects on transmission disequilibrium tests, 565

Eskimos. An HLA class-II allele frequent in ... and Amerindians is found in the Tyrolean Ice Man. 363 Esteban. E.

see Giraldo, M. P. et al.

Evans, K. L.

see Le Hellard, S. et al.

exclusion analyses, LOD score ... for candidate genes using random population samples, 313

Faradz, M. H., Leggo, J., Murray, A., Lam-Po-Tang, P. R. L., Buckley, M. F. & Holden, J. J. A.

distribution of FMR1 and FMR2 alleles in Javanese: confirmation of a specific AGG-interruption pattern in Asian populations, 127

Fae. I

see Fischer, G. F. et al.

Farnham, N. J.

see Camp, N. J. et al.

Fellous, M.

see Khodjet El Khil et al.

Ferguson, P.

see Olivarez, L. et al.

Field, L. L.

see Corder. M. A. et al.

Fischer, G. F., Fae, I., Mann, D., Kriks, D., Jager, W., Platzer, W., Mayr, W. R. & Volc-Platzer, B.

an HLA class-II allele frequent in Eskimos and Amerindians is found in the Tyrolean Ice Man, 363

FMR1 and FMR2. Distribution of ... alleles in Javanese confirmation of a specific

AGG-interruption pattern in Asian populations, 127 Foley, M.

see Underhill, P. A. et al.

Fordyce, A.

see Swerdlow, A. J. et al.

Franceschi, C.

see Varcasia, O. et al.

Franchioso, F

see Martinelli, M. et al.

Fraser, J. S.

see King, A. L. et al.

Fujitani, N.

see Pang. H. et al.

Fulchignoni-Lataud, M. C.

see Greco. L. et al.

Fuster, C

see Cirigliano, V. et al.

Gaggana. M.

see Olivarez, L. et al.

gamete competition model. SNPs and snails and puppy dog's tails: analysis of SNP haplotype data using the 483

gametic disequilibrium. Sampling variance and distribution of the D' measure of overall... between multiallelic loci, 395

Garasto, S.

see Varcasia, O. et al.

Garcia, O.

see Malaspina. P. et al.

Gasparini, P.

see Greco. L. et al.

Gavrila. L.

see Malaspina, P. et al.

Giparaki, M

see Malaspina, P. et al.

Giraldo, M. P., Esteban, E., Aluja, M. P., Nogues, R. M., Backes-Duro, Ch., Dugoujon, M. & Moral, P.

Gm and Km data in two Spanish Pyrenean populations (Andorra and Pallars Sobira): a review of Gm variation in the Mediterranean basin, 537

Greco, L., Babron, M. C., Corazza, G. R., Percopo, S., Sica, R., Clot, F., Fulchignoni-Lataud, M. C., Zavattari, P., Momigliano-Richiardi, P., Casari, G., Gasparini, P., Tosi, R., Mantovani, V., De Virgiliis, S., Lacono, G., D'Alfonso, A., Selinger-Leneman, H., Lemainque, A., Serre, J. L. & Clerget-Darpoux, F.

Existence of a genetic risk factor on chromosome 5q in Italian Coeliac Disease families, 35

glycoprotein (fetuin). Haplotype analysis of human alpha2-HS...gene, 27

Gm and Km data in two Spanish Pyrenean populations (Andorra and Pallars Sobira): review of Gm variation in the Mediterranean basin, 537

Gomez-Casado, E.

see Martinez-Laso, J. et al.

Grover, S. K

see Qadar Pasha, M. A. et al.

Guimaraes, P. E. M.

see Bandelt, H. J. et al.

Gusmao, L.

see Trovoada, M. J. et al.

Gusmao, L., Alves. C. & Amorim, A.

molecular characterisation of four human Y-specific microsatellites (DYS434, DYS437, DYS438, DYS439) for population and forensic studies, 285

haplotype analysis of human alpha2-HS glycoprotein (fetuin) gene, 27

haplotype method. A permutation procedure for the ...
for identification of disease-predisposing variants,
189

Hardy-Weinberg disequilibrium. Fine-scale mapping using, 207

Hata, Y

see Ohashi, J. et al.

- Henke, J.
 - see Osawa, M. et al.
- Hennekam, R. C.M.
 - see Jira. P. E. et al.
- Hermon, C. Alberman, E., Beral, V. & Swerdlow, A. J. mortality and cancer incidence in persons with Down's syndrome, their parents and siblings, 167
- Hermon. C.
 - see Swerdlow, A. J. et al.
- histocompatibility complex. How selection shapes variation of human major, 1
- HLA class-II. An ... allele frequent in Eskimos and Amerindians is found in the Tyrolean Ice Man, 363
- HLA molecular markers in Tuvinians: a population with both Oriental and Caucasoid characteristics, 245
- Holden, J. J. A.
- see Faradz, M. H. et al.
- Hopper, J. L.
- see Visscher, P. M. et al.
- HRAS1 minisatellite. Length variability and interspersion patterns of the ...: a new approach for the reconstruction of human population relationships.
- IDDM11. Grade-of-membership sibpair linkage analysis maps... to chromosome 14q24.3-q31. 387
- Irobi, J., Nelis, E., Meuleman, J., Venken, K., De Jonghe, P., Van Broeckhoven, C. & Timmerman, V
- exclusion of 5 functional candidate genes for distal hereditary motor neuropathy type II (distal HMN II) linked to 12q24.3, 517
- Italian. Existence of a genetic risk factor on chromosome 5q in... Coeliac Disease families. 35
- Italian Veneto speakers. Mitochondrial DNA sequence diversity in two groups of ... from Veneto, 153
- Jacobs, P. A.
 - see Swerdlow, A. J. et al.
- Jager, W
- see Fischer, G. F. et al.
- Jansen, H., Waterwoth, D. M., Nicaud, V., Enhnolm, C., Talmud, P. J., on behalf of the Ears-II-Study Group
- interaction of the common apolipoprotein C-III (APOC3 - 482C > T) and hepatic lipase (LIPC -514C > T) promoter variants affects glucose tolerance in young adults. European Atherosclerosis Research Study II (EARS-II). 237
- Javanese. Distribution of FMR1 and FMR2 alleles in ...: confirmation of a specific AGG-interruption pattern in Asian populations, 127
- Jiang, R., Dong, J. P., Wang, D. & Sun, F. Z.
 - fine-scale mapping using Hardy-Weinberg disequilibrium. 207
- Jira, P. E., Wanders, R. J. A., Smeitink, J. A. M., De Jonk, J. Wevers, R. A., Oosthein, W. Tuerlings, J. H. A. M., Hennekam, R. C. M., Sengers, R. C. A., Waterham, H. R.
 - novel mutations in the 7-dehydrocholesterol reductase gene in 13 patients with Smith-Lemli-Opitz syndrome, 229
- Jeune, B.
 - see Varcasia. O. et al.
- Kanenko, M.
- see Osawa, M. et al.

- Kawasaki, T.
 - see Pang. H. et al.
- Keavney, B.
- see Sinsheimer, J. S. et al.
- Kerimova, G.
- see Malaspina. P. et al.
- Kimura. H.
- see Pang, H. et al.
- King, A. L., Fraser, J. S., Moodie, S. J., Curtis, D., Dearlove, A. M., Ellis, H. J., Rosen-Bronson, S. & Ciclitira, P. J.
- Coeliac Disease: follow-up linkage study provides further support for existence of a susceptibility locus on chromosome 11p11, 377
- Khan, A. P
 - see Qadar Pasha. M. A. et al.
- Khodjet El Khil, H., Triki Marrakchi, R., Yacoubi Loueslati, B., Langaney, A., Fellous, M. & Benammar Elgaaied, A.
- Y chromosome microsatellite variation in three populations of Jerba Island (Tunisia), 263
- Kitano, T.
- see Osawa, M. et al.
- Koda, Y
- see Pang. H. et al.
- Komata, T.
- see Ohashi, J. et al.
- Konkov, V. I.
- see Martinez-Laso, J. et al.
- Kovatchev. D.
- see Malaspina. P. et al.
- Kozlov, A. I.
- see Malaspina, P. et al.
- Kriks, D.
 - see Fischer, G. F. et al.
- Lacono, G.
 - see Greco. L. et al.
- Laluela-Fox, C., Calderon, F. L., Calafell, F., Morera, B. & Bertranpetit. J
- mtDNA from extinct Tainos and the peopling of the Caribbean, 137
- Lam-Po-Tang, P. R. L.
- see Faradz, M. H. et al.
- Langanev, A.
- see Khodjet El Khil et al.
- Lange, K
- see Sinsheimer, J. S. et al.
- Larruga, J. M.
 - see Bandelt. H. J. et al.
- Leggo, J.
- see Faradz. M. H. et al.
- Le Hellard, S., Semple, C. A. M., Morris, D. J. Porteous, D. J. & Evans, K. L.
- physical mapping: integrating computational and molecular genetic data. 221
- Lemainque, A
- see Greco. L. et al.
- Lewin, P.
- see Cirigliano, V. et al.
- see Deng, H. W. et al.
- Li. H.
 - a permutation procedure for the haplotype method for identification of disease-predisposing variants, 189

Lin. A.A

see Underhill. P. A. et al.

Lin, S., Cheng, R. & Wright, F. A.

genetic crossover interference in the human genome, 79

linkage. Family based tests of association and/or, 407 linkage analysis. Grade-of-membership sibpair... maps IDDM11 to chromosome 14q24.3-q31, 387

linkage analysis. A novel method of two-locus ... applied to a genome scan for late onset Alzheimer's disease. 473

linkage analysis of three candidate regions of chromosome f in nonsyndromic familial orofacial cleft, 465

linkage statistics. Sample size requirements to control for stochastic variation in magnitude and location of allele-sharing... in affected sibling pairs, 491

linkage studies. Correcting for multiple analyses in genomewide, 577

linkage study. Coeliac Disease: follow-up... provides further support for existence of a susceptibility locus on chromosome 11p11, 377

lipase. Interaction of the common apolipoprotein C-III (APOC3 -482C > T) and hepatic ... (LIPC -514C > T) promoter variants affects glucose tolerance in young adults. European Atherosclerosis Research Study II (EARS-II), 237

LOD score exclusion analyses for candidate genes using random population samples, 313

longevity. Replication studies in ...: puzzling findings in Danish centenarians at the 3' APOB-VNTR locus, 371

Loutradis, A. see Malaspina, P. et al.

Malaspina, P., Tsopanomichalou, M., Duman, T., Stefan, M., Silvestri, A., Rinaldi, B., Garcia, O., Giparaki, M., Plata, E., Kozlov, A. I., Barbujani, G., Vernesi, C., Papola, F., Ciaverella, G., Kovatchev, D., Kerimova, G., Anagnou, N., Gavrila, L., Veneziano, L., Akar, X. Loutradis, A., Michalodimitrakis, E. X., Terrenato, L. & Novelletto, A.

a multistep process for the dispersal of a Y chromosomal lineage in the Mediterranean area, 339

Macaulay, V. see Pereira, L. et al.

MacPherson, J. N. see Ennis, S. et al.

McKenzie, C. A. C.

see Sinsheimer, J. S. et al.

Malcolm, S. see Prescott, N. J. et al.

Malyarchuk, B. A. & Derenko, M. V.

mitochondrial DXA variability in Russians and Ukrainians: implication to the origin of the Eastern

Slavs, 63 Mann, D. see Fischer, G. F. et al.

Manton, K. G. see Corder, M. A. et al.

Mantovani, V. see Greco, L. et al.

Martinelli, M., Scapoli, L., Pezzetti, F., Carinci, F., Francioso, F., Baciliero, U., Padula, E., Carinci, P., & Tognon, M.

linkage analysis of three candidate regions of chromosome 1 in nonsyndromic familial orofacial cleft, 465 Martinez-Laso, J., Sarkatova, M., Allende, L., Konkov, V. I., Moscoso, J., Silvera-Redondo, C., Pacho, A., Trapaga, J., Gomez-Casado, E. & Arnaiz-Villena, A. HLA molecular markers in Tuvinians: a population

HLA molecular markers in Tuvinians: a population with both Oriental and Caucasoid characteristics, 245

Matsushita, M.

see Ohashi. J. et al.

Mayr. W. R.

see Fischer, G. F. et al.

Mediterranean area. A multistep process for the dispersal
of a Y chromosomal lineage in the, 339

Meyer, D. & Thomson, G.

how selection shapes variation of human major histocompatibility complex. 1

Meuleman, J

see Irobi, J. et al.

Michalodimitrakis, E. N. see Malaspina, P. et al.

Mirazoni Lahr, P. J. see Underhill, P. A. et al.

mitochondrial DXA sequence diversity in two groups of Italian, Veneto speakers from Veneto. 153

mitochondrial DNA variability in Russians and Ukrainians: implication to the origin of the Eastern Slavs, 63

mitochondrial haplogroup L3e. Phylogeography of the human...: a snapshot of African prehistory and Atlantic slave trade, 549

Mogentale-Profizi, N., Chollet, L., Stevanovitch, A., Dubut, V., Poggi, C., Pradie, M. P., Spadoni, J. L. & Beraud-Colomb, E.

mitochondrial DNA sequence diversity in two groups of Italian Veneto speakers from Veneto, 153

Momigliano-Richiardi, P. see Greco, L. et al.

Moodie, S. J. see King, A. L. et al.

see Giraldo, M. P. et al.

Morera, B.

see Laluela-Fox. C. et al. Morris, A. P.

see Whittaker, J. C. Morris, D. J.

Moral, P

see Le Hellard, S. et al.

motor neuropathy type II. Exclusion of 5 functional candidate genes for distal hereditary ... (distal HMN II) linked to 12q24.3, 517

Murray. A.

see Faradz. M. H. et al.

Murray, A.

see Ennis, S. et al.

Morton, N. E.

see Ennis, S. et al.

Moscoso, J.

see Martinez-Laso, J. et al.

Mozambique. Prehistoric and historic traces in the mtDNA of ...: insights into the Bantu expansions and the slave trade. 439

 mtDNA from extinct Tainos and the peopling of the Caribbean, 137

mtDNA. Prehistoric and historic traces in ... of Mozambique: insights into the Bantu expansions and the slave trade, 439

mutation analysis approach. Estimate of the frequency of Wilson's disease in the U.S. Caucasian population: a, 459

Nicaud, V.

see Jansen, H. et al.

Nelis, E,

see Irobi, J. et al.

Nogues, R. M.

see Giraldo, M. P. et al.

Norbo, T.

see Qadar Pasha, M. A. et al.

North, B. V.

see Curtis, D. et al.

Novelletto, A.

see Malaspina, P. et al.

Oefner, P. J.

see Underhill, P. A. et al.

Ogaki, T.

see Pang, H. et al.

Ohashi, J., Yamamoto, S., Tsuuchiya, N., Hata, Y., Komata, T., Matsushita, M., & Tokunaga, K.

comparison of statistical power between 2×2 allele frequency and allele positivity tables in case control studies of complex disease genes, 197

Olivarez, L., Gaggana, M., Pass, K. A., Ferguson, P. & Brewer, G. J.

estimate of the frequency of Wilson's disease in the U.S. Caucasian population: a mutation analysis approach, 459

Oriental. HLA molecular markers in Tuvinians: a population with both ... and Caucasoid characteristics, 245

orofacial cleft. Linkage analysis of three candidate regions of chromosome 1 in nonsyndromic familial, 465

Osawa, M., Yuasa, I., Kitano, T., Henke, J., Kanenko, M., Udono, T., Saitou, N. & Umetsu, K.

haplotype analysis of human alpha2-HS glycoprotein (fetuin) gene. 27

Oosthein, W.

see Jira. P. E. et al.

Pacho, A.

see Martinez-Laso, J. et al.

Padula, E.

see Martinelli, M. et al.

Pallaud, C.

see Tregouet. D. A. et al.

Pang, H., Koda, Y., Soejima, M., Fujitani, X., Ogaki, T., Saito, A., Kawasaki, T. & Kimura, H.

polymorphism of the human ABO-secretor locus (FUT2) in four populations in Asia: indication of distinct Asian subpopulations, 429

Papola. F

see Malaspina, P. et al.

Parkinsonism. Genetics of ...: a review, 111

Pass, K. A.

see Olivarez, L. et al.

Passarino, G.

see Underhill, P. A. et al.

Pena, S. D. J.

see Bandelt, H. J. et al.

Percopo, S.

see Greco. L. et al.

Pereira, L., Macaulay, V. Torroni, A., Scozzari, R., Prata, M. J. & Amorim, A.

prehistoric and historic traces in the mtDNA of Mozambique: insights into the Bantu expansions and the slave trade, 439

Pereira, L.

see Bandelt, H. J. et al.

permutation. A... procedure for the haplotype method for identification of disease-predisposing variants, 189

Pezzetti, F.

see Martinelli, M. et al.

phylogeography. The \dots of Y chromosome binary haplotypes and the origins of modern human populations.

phylogeography of the human mitochondrial haplogroup L3e: a snapshot of African prehistory and Atlantic slave trade, 549

physical mapping: integrating computational and molecular genetic data, 221

lata. E.

see Malaspina, P. et al.

Platzer, W.

see Fischer, G. F. et al.

Poggi, C.

see Mogentale-Profizi, N. et al.

Porteous, D. J.

see Le Hellard, S. et al.

Pradie, M. P.

see Mogentale-Profizi, N. et al.

Prado, V. F

see Bandelt, H. J. et al.

Prata, M. J.

see Bandelt, H. J. et al.

Prata. M. J.

see Trovoada, M. J. et al.

Prata. M. J.

see Pereira, L. et al.

preimplantation genetics: a review, 331

Prescott, N. J., Winter, R. M. & Malcolm, S.

nonsyndromic cleft lip and palate: complex genetics and environmental effects, 505

Qadar Pasha, M. A., Khan, A. P., Ratan Kumar, S. K., Grover, S. K., Ram, R. B., Norbo, T., Srivastava, K. R., Selvamurthy, W. & Brahmachari, S. K.

angiotensin converting enzyme insertion allele in relation to high altitude adaptation, 531

QTL. Power of regression and maximum likelihood methods to map... from sib-pair and DZ twin data, 583

quantitative fluorescent PCR (QF-PCR). Assessment of new markers for the rapid detection of an euploidies by \dots , 421

Ram, R. B.

see Qadar Pasha, M. A. et al.

Ratan Kumar, S. K.

see Qadar Pasha, M. A. et al.

Recker, R. R.

see Deng, H. W. et al.

regression. Power of ... and maximum likelihood methods to map QTL from sib-pair and DZ twin data, 583

Rengo, C. see Bandelt, H. J. et al.

Rinaldi B

see Malaspina. P. et al.

Rizza, T.

see Varcasia, O. et al.

Rodriguez, S

see Zapata, C. et al.

Rosen-Bronson, S.

see King, A. L. et al.

Russians and Ukranians. Mitochondrial DNA variability in . . . : implication to the origin of the Eastern Slavs, 63

Saito, A.

see Pang, H. et al.

Saitou, N

see Osawa, M. et al.

Salas, A.

see Vega, A. et al.

Santos, M. S.

see Bandelt, H. J. et al.

São Tomé e Principé. Evidence for population substructuring in ... as inferred from Y-chromosome STR analysis, 271

Sarkatova, M.

see Martinez-Laso, J. et al.

Sass. C

see Tregouet, D. A. et al.

Scapoli, L.

see Martinelli, M. et al.

Scozzari, R.

see Bandelt, H. J. et al.

Scozzari, R.

see Pereira, L. et al.

Selinger-Leneman, H.

see Greco, L. et al.

Selvamurthy, W

see Qadar Pasha, M. A. et al.

Semple, C. A. M.

see Le Hellard, S. et al.

Sengers, R. C. A.

see Jira, P. E. et al.

Serre, J. L.

see Greco, L. et al.

sex chromosome abnormalities. Mortality and cancer incidence in persons with numerical...: a cohort study, 177

Sham, P. C.

see Curtis, D. et al.

Shen, R. A.

see Underhill, P. A. et al.

sib-pair. Power of regression and maximum likelihood methods to map QTL from ... and DZ twin data, 583

Sica. R.

see Greco, L. et al.

Silvera-Redondo, C.

see Martinez-Laso, J. et al.

Silvestri, A.

see Malaspina, P. et al.

Sinsheimer, J. S., McKenzie, C. A. C., Keavney, B. & Lange, K.

SNPs and snails and puppy dog's tails: analysis of SNP haplotype data using the gamete competition model, 483

slave trade. Phylogeography of the human mitochondrial haplogroup L3e: a snapshot of African prehistory and Atlantic, 549

slave trade. Prehistoric and historic traces in the mtDNA of Mozambique: insights into the Bantu expansions and the, 439

Smeitink, J. A. M.

see Jira, P. E. et al.

Smith-Lemli-Opitz syndrome. Novel mutations in the 7-dehydrocholesterol reductase gene in 13 patients with, 229

SNPs and snails and puppy dog's tails: analysis of SNP haplotype data using the gamete competition model, 483

Soejima, M

see Pang. H. et al.

Spadoni, J. L.

see Mogentale-Profizi, N. et al.

Spanish Pyrenean populations, Gm and Km data in two ... (Andorra and Pallars Sobira): a review of Gm variation in the Mediterranean basin, 537

Srivastava, K. R.

see Qadar Pasha, M. A. et al.

Stefan, M.

see Malaspina, P. et al.

Stevanovitch, A

see Mogentale-Profizi, N. et al.

stochastic variation. Sample size requirements to control for ... in magnitude and location of allele-sharing linkage statistics in affected sibling pairs, 491

Sun. F. Z.

see Jiang, R. et al.

Szpiro-Tapies, S

see Cirigliano, V. et al.

Swerdlow, A. J.

see Hermon, C. et al.

Swerdlow, A. J., Hermon, C., Jacobs, P. A., Alberman, E., Beral, V., Daker, M., Fordyce, A. & Youings, S. mortality and cancer incidence in persons with numerical sex chromosome abnormalities: a cohort

Talmud, P. J.

see Jansen, H. et al.

study, 177

Tan. Q.

see Varcasia, O. et al.

Tapper, W

see Ennis, S. et al.

TDT-type methods. Sample size calculations for classical association and \dots using family data, 293

Terrenato. L.

see Malaspina, P. et al.

Thomson, G.

see Meyer, D.

Timmerman, V.

see Irobi, J. et al.

Tiret, L.

see Tregouet, D. A. et al.

Tognon, M.

see Martinelli, M. et al.

Tokunaga, K.

see Ohashi, J. et al.

Torroni, A.

see Bandelt, H. J. et al.

Torroni, A.

see Pereira. L. et al.

Tosi R

see Greco. L. et al.

transmission disequilibrium test. Epistasis and its possible effects on, 565

Trapaga, J.

see Martinez-Laso, J. et al.

Tregouet, D. A., Pallaud, C., Sass, C., Visvikis, S. & Tiret, L.

sample size calculations for classical association and TDT-type methods using family data, 293

Triki Marrakchi, R.

see Khodjet El Khil et al.

Trovoada, M. J., Alves, C., Gusmao, L., Abade, A., Amorim, A. & Prata, M. J.

evidence for population sub-structuring in São Tomé e Principé as inferred from Y-chromosome STR analysis, 271

Tsopanomichalou. M.

see Malaspina, P. et al.

Tsuuchiya, N.

see Ohashi, J. et al.

Tuerlings, J. H. A. M. see Jira, P. E. et al.

Tunisia. Y chromosome microsatellite variation in three populations of Jerba Island. 263

Tunivians. HLA molecular markers in ...: a population with both Oriental and Caucasoid characteristics, 245

Tyrolean Ice Man. An HLA class-II allele frequent in Eskimos and Amerindians is found in the, 363

Udono, T

see Osawa, M. et al.

Umetsu, K.

see Osawa, M. et al.

Underhill, P. A., Passarino, G., Lin, A. A., Shen, R. A., Foley, M., Mirazoni Lahr, P. J., Oefner, P. J. & Cavalli-Sforza, L. L.

the phylogeography of Y chromosome binary haplotypes and the origins of modern human populations,

Van Broeckhoven, C.

see Irobi, J. et al.

Varcasia, O., Garasto, S., Rizza, T., Andersen-Ranberg, K., Jeune, B., Bathum, L., Andreev, K., Tan, Q., Yashin, A. I., Bonafe, M., Franceschi, C. & De Benedictis, G.

replication studies in longevity: puzzling findings in Danish centenarians at the 3' APOB-VNTR locus, 371

Vaughan, J. R., Davis, M. B. & Wood, N. W genetics of Parkinsonism: a review, 111

Vega, A., Salas, A., Costas, J., Barros, F. & Carracedo, A. length variability and interspersion patterns of the HRAS1 minisatellite: a new approach for the reconstruction of human population relationships, 351

Veneziano, L.

see Malaspina, P. et al.

Venken, K.

see Irobi, J. et al.

Vernesi, C.

see Malaspina, P. et al.

Visscher, P. M. & Hopper, J. L.

power of regression and maximum likelihood methods to map QTL from sib-pair and DZ twin data, 583 Visvikis, S.

see Tregouet, D. A. et al.

Volc-Platzer, B.

see Fischer, G. F. et al.

Wanders, R. J. A.

see Jira. P. E. et al.

Wang. D.

see Jiang. R. et al.

Waterham, H. R.

see Jira. P. E. et al.

Waterwoth, D. M.

see Jansen, H. et al.

Wevers, R. A.

see Jira. P. E. et al.

Whittaker, J. C. & Morris, A. P.

family based tests of association and/or linkage, 407 Wilson, S. R.

epistasis and its possible effects on transmission disequilibrium tests, 565

Wilson's disease. Estimate of the frequency of ... in the U. S. Caucasian population: a mutation analysis approach, 459

Winter, R. M.

see Prescott, N. J. et al.

Wood, N. W.

see Vaughan, J. R. et al.

Woodbury, M. A.

see Corder, M. A. et al.

Wright, F. A. see Lin, S. et al.

200 200 000

Yacoubi Loueslati, B. see Khodjet El Khil et al.

Yamamoto, S.

see Ohashi, J. et al.

Yashin, A. I.

see Varcasia, O. et al.

Y chromosomal. A multistep process for the dispersal of a ... lineage in the Mediterranean area, 339

Y chromosome. The phylogeography of ... binary haplotypes and the origins of modern human populations, 43

Y chromosome microsatellite variation in three populations of Jerba Island (Tunisia), 263

Y-chromosome STR. Evidence for population substructuring in São Tomé e Principé as inferred from ... analysis, 271

Y-specific microsatellites. Molecular characterisation of four human...(DYS434, DYS437, DYS438, DYS-439) for population and forensic studies, 285

Yuasa, I.

see Osawa, M. et al.

Zapata, C., Carollo, C. & Rodriguez, S.

sampling variance and distribution of the D' measure of overall gametic disequilibrium between multiallelic loci, 395

Zavattari. P

see Greco, L. et al.

12q24.3. Exclusion of 5 functional candidate genes for distal hereditary motor neuropathy type II (distal HMN II) linked to, 517

7-dehydrocholesterol reductase gene. Novel mutations in the ... in 13 patients with Smith-Lemli-Opitz syndrome, 229

